

ENTERED

OIPE

RAW SEQUENCE LISTING DATE: 03/07/2002 PATENT APPLICATION: US/10/081,126 TIME: 10:01:17

Input Set : A:\AR4951.txt

```
4 <110> APPLICANT: De Vries, Gerald W.
      6 <120> TITLE OF INVENTION: Methods of Extending Corneal Graft
              Survival
      9 <130> FILE REFERENCE: P-AR 4951
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/081,126
C--> 11 <141> CURRENT FILING DATE: 2002-02-22
     11 <160> NUMBER OF SEQ ID NOS: 7
     13 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     15 <210> SEQ ID NO: 1
     16 <211> LENGTH: 4113
     17 <212> TYPE: DNA
     18 <213> ORGANISM: Homo sapiens
     20 <220> FEATURE:
     21 <221> NAME/KEY: CDS
     22 <222> LOCATION: (22)...(4110)
     24 <400> SEQUENCE: 1
     25 acccaegege ageggeegga g atg eag egg gge gee geg etg tge etg ega
                                Met Gln Arg Gly Ala Ala Leu Cys Leu Arg
     27
                                                                           99
     29 ctg tgg ctc tgc ctg gga ctc ctg gac ggc ctg gtg agt gac tac tcc
     30 Leu Trp Leu Cys Leu Gly Leu Leu Asp Gly Leu Val Ser Asp Tyr Ser
                         15
                                             20
     33 atg acc ccc ccg acc ttg aac atc acg gag gag tca cac gtc atc gac
     34 Met Thr Pro Pro Thr Leu Asn Ile Thr Glu Glu Ser His Val Ile Asp
                     30
    35
                                         35
     37 acc ggt gac agc ctg tcc atc tcc tgc agg gga cag cac ccc ctc qag
                                                                           195
     38 Thr Gly Asp Ser Leu Ser Ile Ser Cys Arg Gly Gln His Pro Leu Glu
                                     50
     41 tgg get tgg eea gga get eag gag geg eea gee ace gga gae aag gae
                                                                           243
     42 Trp Ala Trp Pro Gly Ala Gln Glu Ala Pro Ala Thr Gly Asp Lys Asp
                                 65
    45 age gag gac acg ggg gtg gtg cga gac tgc gag ggc aca gac gcc agg
                                                                           291
    46 Ser Glu Asp Thr Gly Val Val Arg Asp Cys Glu Gly Thr Asp Ala Arg
    47 75
                             80
    49 ccc tac tgc aag gtg ttg ctg ctg cac gag gta cat gcc aac gac aca
                                                                           339
     50 Pro Tyr Cys Lys Val Leu Leu Leu His Glu Val His Ala Asn Asp Thr
                                            100
     53 ggc agc tac gtc tgc tac tac aag tac atc aag gca cgc atc gag ggc
                                                                           387
     54 Gly Ser Tyr Val Cys Tyr Tyr Lys Tyr Ile Lys Ala Arg Ile Glu Gly
    55
                    110
                                        115
     57 acc acg gcc gcc agc tcc tac gtg ttc gtg aga gac ttt gag cag cca
                                                                           435
    58 Thr Thr Ala Ala Ser Ser Tyr Val Phe Val Arg Asp Phe Glu Gln Pro
    59
                125
```

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61 ttc atc aac								483
62 Phe Ile Ası	n Lys Pro	Asp Thr	Leu Leu	Val Asn	Arg Lys	Asp Ala	Met	
63 140		145			150			
65 tgg gtg ccd								531
66 Trp Val Pro	Cys Leu	Val Ser	Ile Pro	Gly Leu	Asn Val	Thr Leu	Arg	
67 155		160		165			170	
69 tcg caa ago								579
70 Ser Gln Ser	Ser Val	Leu Trp	Pro Asp	Gly Gln	Glu Val	Val Trp	Asp	
71	175			180		185		
73 gac cgg cgg			_	_	-		-	627
74 Asp Arg Arg	g Gly Met	Leu Val	Ser Thr	Pro Leu	Leu His	Asp Ala	Leu	
75	190		195			200		
77 tac ctg cag	g tgc gag	acc acc	tgg gga	gac cag	gac ttc	ctt tcc	aac	675
78 Tyr Leu Glr	n Cys Glu	Thr Thr	Trp Gly	Asp Gln	Asp Phe	Leu Ser	Asn	
79 205	5		210		215			
81 ccc ttc ctg								723
82 Pro Phe Let	ı Val His	Ile Thr	Gly Asn	Glu Leu	Tyr Asp	Ile Gln	Leu	
83 220		225			230			
85 ttg ccc agg	g aag tcg	ctg gag	ctg ctg	gta ggg	gag aag	ctg gtc	ctc	771
86 Leu Pro Arg	J Lys Ser	Leu Glu	Leu Leu	Val Gly	Glu Lys	Leu Val	Leu	
87 235		240		245			250	
89 aac tgc acc								819
90 Asn Cys Thi	. Val Trp	Ala Glu	Phe Asn	Ser Gly	Val Thr	Phe Asp	Trp	
91	255			260		265		
93 gac tac cca								867
94 Asp Tyr Pro		Gln Ala	Glu Arg	Gly Lys	Trp Val	Pro Glu	Arg	
95	270		275			280		
97 cgc tcc caa	_		_	-	_			915
98 Arg Ser Glr		His Thr		Ser Ser		Thr Ile	His	
99 285			290		295			
101 aac gtc ag						-		963
102 Asn Val Se	er Gln Hi	_	-	r Tyr Val		s Ala Asn	Asn	
103 300		305			310			
105 ggc atc ca								1011
106 Gly Ile Gl	n Arg Ph		i Ser Th			. His Glu		
107 315		320		325			330	
109 ccc ttc at								1059
110 Pro Phe II			b Leu Ly		o lle Leu			
111	33			340		345		3 4 0 5
113 gca gga ga								1107
114 Ala Gly As	_	u vai Lys		_	s Leu Ala		Pro	
115	350		35			360		
117 ccg ccc ga								1155
118 Pro Pro Gl		n Trp Tyl		о ста газ		_	Arg	
119 36			370		375			1000
121 cac agt co	a cat go	c ctg gtg	ctc aag	g gag gtg	g aca gag	gec age	aca	1203
122 His Ser Pr	O HIS AL			s Glu Val		ara ser	rnr	
123 380	±	385			390			1051
125 ggc acc ta	ic acc ct	u gad atg	, tgg aad	s too got	r der dde	ctg agg	cgc	1251

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Input Set : A:\AR4951.txt

	Gly 395	Thr	Tyr	Thr	Leu	Ala 400	Leu	Trp	Asn	Ser	Ala 405	Ala	Gly	Leu	Arg	Arg 410	
129	aac	atc	agc	cta	σασ	cta	at.a	at.a	aat.	at.a	ccc	ccc	caq	ata	cat	σασ	1299
		Ile															
	HDII	110	501	Dea		neu	vul	Vul	A5II		rio	FIO	GIII	116		GIU	
131					415			_	_	420					425		
		gag															1347
134	Lys	Glu	Ala	Ser	Ser	Pro	Ser	Ile	Tyr	Ser	Arg	His	Ser	Arg	Gln	Ala	
135				430					435					440			
137	ct.c	acc	tac	acσ	acc	tac	aaa	at.a	ccc	cta	cct	ctc	aαc	atc	cag	taa	1395
		Thr															1000
139	пси		445	1111	niu	- y	OLY	450	110	Leu	110	пеа	455	116	GIII	11P	
		.															4 4 4 4
		tgg															1443
	His	\mathtt{Trp}	Arg	Pro	\mathtt{Trp}	Thr	Pro	Cys	Lys	Met	Phe	Ala	Gln	Arg	Ser	Leu	
143		460					465					470					
145	cgg	cgg	cgg	cag	cag	caa	gac	ctc	atg	cca	cag	tgc	cgt	gac	tgg	agg	1491
		Arg															
147		,	,			480	-				485	_				490	
		gtg	200	200	asa		~~~	ata	220	000	-	~~~	3.00	ata	~~~		1539
																	1339
	Ата	Val	TIIL	THE		ASP	Ата	vai	ASII		тте	GIU	ser	Leu	-	THE	
151					495					500					505		
		acc															1587
154	Trp	Thr	Glu	Phe	Val	Glu	Gly	Lys	Asn	Lys	Thr	Val	Ser	Lys	Leu	Val	
155				510					515					520			
157	atc	cag	aat	gcc	aac	qtq	tct	qcc	atq	tac	aaq	tqt	qtq	qtc	tcc	aac	1635
		Gln															
159			525			,		530		-1-		0,10	535		501	11011	
	224	gtg		~~~	ant.	~~~	~~~	_	2 t a	+	++~	+-+		200		n+ n	1602
																	1683
	гĀЗ	Val	GIY	GII	Asp	GIU	_	Leu	тте	Tyr	Pne	_	val	Thr	Thr	TTE	
163		540					545					550					
		gac															1731
166	Pro	Asp	Gly	Phe	Thr	Ile	Glu	Ser	Lys	Pro	Ser	Glu	Glu	Leu	Leu	Glu	
167	555					560					565					570	
169	qqc	cag	ccq	ata	ctc	cta	agc	tac	caa	acc	gac	agc	tac	aaq	tac	σασ	1779
		Gln															
171	0-1	V			575		201	0,0	0	580	p	001	- 1 -	~75	585	014	
	aa+	ata	000	+~~		000	ata	226	a+ ~		200	a+~	~~~	~~+		~~~	1007
		ctg															1827
	HIS	Leu	Arg		Tyr	Arg	ren	Asn		ser	Thr	Leu	HIS	_	Ala	HIS	
175				590					595					600			
177	ggg	aac	ccg	ctt	ctg	ctc	gac	tgc	aag	aac	gtg	cat	ctg	ttc	gcc	acc	1875
178	Gly	Asn	Pro	Leu	Leu	Leu	Asp	Cys	Lys	Asn	Val	His	Leu	Phe	Ala	Thr	
179			605				_	610	_				615				
181	cct	ctg	acc	acc	age	cta	gag	gag	at.a	gca	cct	aaa	aca	cac	cac	acc	1923
182	Pro	Leu	Δla	Δla	Ser	Len	Glu	Glu	Val	712	Dro	C117	712	Ara	Uic	712	1710
183		620		43.4.U	JGI	ست	625	JIU	101	AIG	110	630	пта	лту	TTD	NIG	
									- 4-								1074
		ctc															1971
		Leu	ser	Leu	Ser		Pro	Arg	Val	Ala		Glu	His	Glu	Gly		
187						640					645					650	
		gtg															2019
		Val															
	-		-				-	-	-			-	-		-		

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Input Set : A:\AR4951.txt
Output Set: N:\CRF3\03072002\J081126.raw

101					CEE												
191			+	~+~	655	~+~	~~~	~~~	~+~	660	~~~			a+ a	665	~~~	2067
	aag																2067
	Lys	гĀг	TÄT		ser	Val	GIII	Ата		GIU	ATa	PLO	Arg		THE	GIII	
195				670	~+~	~+~	~+~		675			.		680			2115
	aac	-		_		_				_	-	-	_		_	_	2115
	Asn	ьeu		ASP	ьeu	ьец	vaı		vaı	ser	ASP	ser		GIU	мет	GIN	
199	4		685					690					695				0160
	tgc																2163
	Cys		vaı	Ата	GIY	Ala		Ala	Pro	Ser	ITe		Trp	Tyr	Lys	Asp	
203		700					705					710					
	gag																2211
	Glu	Arg	Leu	Leu	Glu		Lys	Ser	GLY	Val	_	Leu	Ala	Asp	Ser		
	715					720					725					730	
	cag																2259
	Gln	Lys	Leu	Ser		Gln	Arg	Val	Arg		GLu	Asp	Ala	Gly		Tyr	
211					735					740					745		
	ctg																2307
	Leu	Cys	Ser		Cys	Arg	Pro	Lys	_	Cys	Val	Asn	Ser		Ala	Ser	
215				750					755					760			
	gtg																2355
	Val	Ala		Glu	Gly	Ser	Glu	Asp	Lys	Gly	Ser	Met	Glu	Ile	Val	Ile	
219			765					770					775				
	ctt																2403
	Leu		Gly	Thr	Gly	Val		Ala	Val	Phe	Phe	_	Val	Leu	Leu	Leu	
223		780					785					790					
	ctc																2451
	Leu	Ile	Phe	Cys	Asn	Met	Arg	Arg	Pro	Ala	His	Ala	Asp	Ile	Lys	Thr	
	795					800					805					810	
	ggc		_				_	_						-			2499
	Gly	\mathtt{Tyr}	Leu	Ser	Ile	Ile	Met	Asp	Pro	Gly	Glu	Val	Pro	Leu	Glu	Glu	
231			·		815					820					825		
	caa																2547
	Gln	Cys	Glu	Tyr	Leu	Ser	Tyr	Asp	Ala	Ser	Gln	${\tt Trp}$	Glu	Phe	Pro	Arg	
235				830					835					840			
	gag		_		_		_						-			-	2595
	Glu	Arg		His	Leu	Gly	Arg	Val	Leu	Gly	Tyr	Gly	Ala	Phe	Gly	Lys	
239			845					850					855				
	gtg																2643
	Val		Glu	Ala	Ser	Ala		Gly	Ile	His	Lys	_	Ser	Ser	Cys	Asp	
243		860					865					870					
	acc		-	_		_	_				_	-	_	_		_	2691
	Thr	Val	Ala	Val	Lys	Met	Leu	Lys	Glu	Gly	Ala	Thr	Ala	Ser	Glu		
	875					880					885					890	
	cgc																2739
	Arg	Ala	Leu	Met		G1u	Leu	Lys	Ile		Ile	His	Ile	Gly		His	
251					895					900					905		
	ctc																2787
	Leu	Asn	Val		Asn	Leu	Leu	Gly		Cys	Thr	Lys	Pro		Gly	Pro	
255				910					915					920			

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						gag											2835
	Leu	Met		Ile	Val	Glu	Phe		Lys	Tyr	Gly	Asn		Ser	Asn	Phe	
259 261	cta	cac	925	aan	caa	gac	acc	930	200	000	tac	aca	935	224	tot	000	2883
						Asp											2003
263		940		-1-	3		945				010	950	0	_10	001		
265	gag	cag	cgc	gga	cgc	ttc	cgc	gcc	atg	gtg	gag	ctc	gcc	agg	ctg	gat	2931
						Phe											
• •	955					960					965					970	
						agc											2979
	Arg	Arg	Arg	Pro	_	Ser	Ser	Asp	Arg		Leu	Phe	Ala	Arg		Ser	
271	aan	300	nan	aaa	975	gcg	3 a a	oaa	aat	980	003	~ 2.0	<i>a</i>	422	985	~~~	3027
						Ala											3027
275	270		014	990	_	7114	**** 9	*****9	99!		110	пор	0111	1000		Gru	
277	gac	ctg	tgg	ctg	agc	ccg	ctg	acc	atg	gaa	gat	ctt	gtc			agc	3075
						Pro											
279			1005					1010					1015				
						ggg											3123
	Phe			Ala	Arg	Gly			Phe	Leu	Ala		-	Lys	Cys	Ile	
283	020	1020		ata	aat	gct	1025		a++	at a	ata	1030		200	~~~	~+~	2171
						Ala											3171
	1035			Deu	1114	1040		21011	110	Deu	1045		Olu	UCI	пър	1050	
			atc	tqt	gac	ttt	-	ctt	qcc	caa	-		tac	aaa	gac		3219
						Phe											
291					1055					1060					1065		
						ggc											3267
	Asp	Tyr	Val			Gly	Ser	Ala	_		Pro	Leu	Lys	_		Ala	
295	aat		300	1070		gac	220	~+~	1075		200	~~~	204	1080		+~~	2215
						Asp											3315
299			1085				212	1090	_			0111	1095	_	, 44	110	
	tcc	ttt	ggg	gtg	ctt	ctc	tgg	gag	atc	ttc	tct	ctg			tcc	ccg	3363
302	Ser	Phe	Gly	Val	Leu	Leu	Trp	Glu	Ile	Phe	Ser	Leu	Gly	Ala	Ser	Pro	
303		1100					1105					1110					
						atc											3411
			GLY	Val	GIn	Ile		Glu	GLu	Phe	_		Arg	Val	Arg	-	
	1115		ann	ato	200	1120		a a a	ata	a cc	1125		000	2+2	aaa	1130	2450
						gcc Ala											3459
311	0 -1				1135			014	DC4	1140		110	1124	110	1145		
313	atc	atg	ctg	aac		tgg	tcc	gga	gac			gcq	aga	cct			3507
						Trp											
315				1150					1155					1160			
						atc											3555
210			T	77-7	C1	T1 ^	Lau	C137	Acn	T.em	T.011	Gln	Glv	Ara	Clv	Τ.Δ11	
	Ser	Asp			GIU	TTE	пец	_	_	LCu	Dea	0111	_	_	GLY	LCu	
319			1165	5		gtc		1170)				1175	5	_		3603

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 03/07/2002 PATENT APPLICATION: US/10/081,126 TIME: 10:01:18

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; Xaa Pos. 2,3,4,5,6,7,8,9,10,11,13

Seq#:6; N Pos. 1,2,4,8,9,10,11,12,13

Seq#:7; N Pos. 1

VERIFICATION SUMMARY

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L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:751 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!

L:755 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:5

L:756 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0

L:768 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!

L:772 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:6

L:773 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0

L:783 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!

L:787 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:7

L:788 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0